#### TRANSMITTAL, LETTER TO THE UNITED STATES DESIGNATED / ELECTED OFFICE (DO/EO/US) **CONCERNING A FILING UNDER 35 U.S.C. 371**

NAL APPLICATION NO PCT/EP00/09241

INTERNATIONAL FILING DATE

21 September 2000

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PRIORITY DATE CLAIMED

21 September 1999

TITLE	CF	INVENTION

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### PEPTIDES AGAINST AUTO-ANTIBODIES CAUSING DCM

Wolfgang ROENSPECK, Rudolf KUNZE, Gerd WALLUKAT -and- Manuela DIERENFELD

Applicant herein submits to the United States Designated/Elected Office (DO/EO/US) the following
items and other information.
This is a FIRST submission of items concerning a filing under 35 U.S.C. 371.
<ol> <li>This is a FIRST submission of items concerning a filing under 35 U.S.C. 371.</li> <li>This is a SECOND or SUBSEQUENT submission of items concerning a filing under 35 U.S.C. 371.</li> </ol>
3. This express request to begin national examination procedures (35 U.S.C. 371(f)) at any time rather than delay examination until the expiration of the applicable time limit set in 35 U.S.C. 371(b) and PCT Articles 22 and 39(1).
4. A proper Demand for Internatl. Preliminary Examination was made by the 19th month from earliest claimed priority date.
5. A copy of the International Application as filed (35 U.S.C. 371(c)(2))
a. $\square$ is transmitted herewith (required only if not transmitted by the International Bureau).
b. has been transmitted by the International Bureau.
c. I is not required, as the application was filed in the United States Receiving Office (RO/US)
6. A translation of the International Application into English (35 U.S.C. 371(c)(2)).
7. Amendments to the claims of the International Application under PCT Article 19 (35 U.S.C. 371(c)(3))
a. are transmitted herewith (required only if not transmitted by the International Bureau).
b.  have been transmitted by the International Bureau.
c. have not been made; however, the time limit for making such amendments has NOT expired.
d. have not been made and will not be made.
8. A translation of the amendments to the claims under PCT Article 19 (35 U.S.C. 371(c)(3)).
9. An oath or declaration of the inventor(s) (35 U.S.C. 371(c)(4)).
10. A translation of the annexes to the Internatl. Preliminary Examination report under PCT Article 36 (35 U.S.C. 371(c)(5)).
Items 11. to 16. below concern other document(s) or information included:
11. An Information Disclosure Statement under 37 CFR 1.97 and 1.98.
12. An assignment document for recording. A separate cover sheet compliance with 37 CFR 3.28 and 3.31 is included.
13. A FIRST preliminary amendment.
A SECOND or SUBSEQUENT preliminary amendment.
14. \( \sigma\) A substitute specification.
15. A change of power of attorney and/or address letter.
16. Other items or information:
International Search Report – EPO
PCT/IB/301 Form
PCT/IB/304 Form
PCT/IB/308 Form First Page of Publication
International Preliminary Examination Report – No Annexes

US APPLICATION NO (If known, see 37 CFR 1.5)	000609	PCT/EP00	l l	P67710	uso l
<u> </u>	088681	r C I/LP U	00241	CALCULATIONS	PTO USE ONLY
17. The following fees	are submitted:				
Basic National Fee (37 (					
Internati. prelim. examina		O (37 CFR 1.492 (a) (	1)) \$710.00		
No international prelimina (a) (2)) but international s	arv examination fee pai	id to USPTO (37 CFR	1.492		
Neither international preli	minary examination fe	e (37 CFR 1.492 (a) (	3))		
International preliminary ( (a) (4)) and all claims sati	examination fee paid to	USPTO (37 CFR 1.4	92		
Search Report prepared					
		OPRIATE BASIC FI		\$ 890.00	
Surcharge of \$130.00 for	furnishing the oath or	declaration later that	n	\$ 130.00	
20 30 months fro	om the earliest claimed	priority date (37 CFR	1.492(e)).		
Claims	Number Filed	Number Extra	Rate		
Total Claims	<b>12 -</b> 20 =	-0-	x \$18.00	\$	
Independent Claims	1 - 3 =	-0-	x \$84.00	\$	
Multiple Dependent Clair	n(s) (if applicable)		+ \$280.00	\$	
	TOTA	L OF ABOVE CAL	CULATIONS =	\$ 1020.00	
Reduction by 1/2 for filing Entity statement must als	g by <b>small entity</b> , if ap so be filed. (Note 37 C	pplicable. Verified Sm FR 1.9, 1.27, 1.28).	all	\$	
			SUBTOTAL =	\$ 1020.00	
Processing fee of \$130 f	Processing fee of \$130 for furnishing the <b>English translation</b> later than 20  30 months from the earliest claimed priority date (37 CFR 1.492(f))				
			TIONAL FEE =	\$ 1020.00	
Fee of \$40.00 for recording the enclosed <b>assignment</b> (37 CFR 1.21(h)). Assignment must be accompanied by appropriate cover sheet (37 CFR 3.28, 3.31).					
	TOTAL FEES ENCLOSED =				
				Amt. to be refunded:	\$
	Amt. charged:	\$			
a. A check in the amount of \$ 1020.00 to cover the above fees is enclosed.  b. Please charge my Deposit Account No. 06-1358 in the amount of \$ to cover the above fees.  A duplicate copy of this sheet is enclosed.  c. The Commissioner is hereby authorized to charge my account any additional fees set forth in §1.492 during the					
pendency of this application, or credit any overpayment to Deposit Account No. <u>06-1358</u> . A duplicate copy of this sheet is enclosed.					
SEND ALL CORRESPONDENCE TO:				M2 - CV	losa
JACOBSON HOLMAN PLLC  400 7th Street, N.W., Suite 600  Wi				iam E. Player	Swh-

400 7th Street, N.W., Suite 600 Washington, DC 20004 202-638-6666

**CUSTOMER NUMBER: 00136** 

William E. Player Reg. No. 31,409

JPH&S 3/95

## IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

Applicants:

Wolfgang ROENSPECK et al

Serial No.:

New

Filing Date:

March 21, 2002

For:

PEPTIDES AGAINST AUTO-ANTIBODIES CAUSING DCM

### **PRELIMINARY AMENDMENT**

Assistant Commissioner of Patents Washington, D.C. 20231

Sir:

Prior to initial examination, please amend the above-identified application as follows:

#### **IN THE SPECIFICATION**

Please insert the following sentence on line 1, immediately following the title:

--This is a 371 of PCT/EP00/09241, filed September 21, 2000, the disclosure of which is incorporated herein by reference.--

#### **IN THE CLAIMS**

Please cancel original claims 1-12 without prejudice or disclaimer.

Please add new claims 13-24 as found on the following pages.

- 2 -

#### CLAIMS:

#### 13. A peptide having the amino acid sequence

X01-X02-X03-G-X04-X05-X06-X07-X08-X09-W-X10-X11-X12

#### wherein

X01 = amino group, acetyl group, biotin group, fluorescent label, spacer, linker or deletion;

X02 = D,G,E,T,S or deletion;

X03 = W,Y,F,G,T;

 $X04 = T_1S_1A_1G_2$ 

X05 = L,F,Y,W;

X06 = V,I,W,F,Y;

X07 = S,A,C;

X08 = G,D,E,N,Q;

X09 = F,L,I,Y;

X10 = E,Q,T,S,L;

X11 = Y,F,T,S,W;

X12 = amide, the free acid, GKK, or a spacer;

and peptides having the amino acid sequence

X01-X02-W-X03-R-X04-X05-X06-X07-X08-E-A-R-X09-X10-X11-X12-X13-X14-X15-X16-X17

#### wherein

X01 = amino group, amino acid, peptide, acetyl group, biotin group,

fluorescent label, spacer, linker or deletion;

14. The peptide according to claim 13 having the following amino acid sequence:

X17 = amide, the free acid, GKK, SGKK or a spacer.

X01-X02-X03-G-X04-X05-X06-X07-X08-X09-W-X10-X11-X12

wherein

X01 = amino group, acetyl group, biotin group, fluorescent label, spacer, linker or deletion;

X02 = D,E,T, or deletion;

X03 = W,Y,T;

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X04 = T,S;
X05 = L,F;
X06 = V,F;
X07 = S;
X08 = G,D,E;
X09 = F,L;
X10 = E,Q,T,L;
X11 = Y,T,S;
X12 = amid, the free acid, GKK, or a spacer;
and peptides having the amino acid sequence
X01-H-W-X03-R-A-X05-S-D-X08-E-A-R-R-S-Y-X12-D-P-X15-X16-X17
wherein
X01 = amino group, amino acid, peptide, acetyl group, biotin group,
fluorescent label, spacer, linker or deletion;
X03 = Y, W;
X05 = T, E;
X08 = G, or a deletion;
X12 = A, N;
X15 = K, T;
X16 \approx S, or a deletion;
X17 = amide, the free acid, GKK, SGKK or a spacer.
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15. The peptides according to claim 13 selected from the group consisting of:

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-TGSFFSELWTSR<sup>2</sup>,
EYGSFFSELWTSR<sup>2</sup>,
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1-5-

TYGTLFSDFWLSR<sup>2</sup>, DWGTLVSGFWEYR<sup>2</sup>, DWGTLFSDFWQTR<sup>2</sup>,

wherein  $R^2$  is an acid amide, a free acid or  $GKKR^3$ , and wherein  $R^3$  is an acid amide or a free acid;

with the proviso that a maximum of one non-conservative amino acid exchange is effected per amino acid position in the sequence, wherein "non-conservative exchange" means an exchange of amino acids between the groups mentioned below:

Group I: Leu, Ile, Val, Met, His, Trp, Tyr, Phe,

Group II: Glu, Gln, Asp, Asn,

Group III: Ser, Thr, Cys, Gly, Ala, Pro,

Group IV: Lys, Arg;

and

peptides selected from the group consisting of:

HWWRAESD-EARRSYNDPK-R<sup>2</sup>, HWYRATSDGEARRSYADPTSR<sup>2</sup>,

with the proviso that a maximum of two non-conservative amino acid exchanges are effected per amino acid position in the sequence, wherein "non-conservative exchange" means an exchange of amino acids between the groups mentioned below:

Group I: Leu, Ile, Val, Met, His, Trp, Tyr, Phe,

Group II: Glu, Gln, Asp, Asn,

Group III: Ser, Thr, Cys, Gly, Ala, Pro,

Group IV: Lys, Arg.

TYGTLFSDFWLSR<sup>2</sup>, DWGTLVSGFWEYR<sup>2</sup>, DWGTLFSDFWQTR<sup>2</sup>,

wherein  $R^2$  is an acid amide, a free acid or  $GKKR^3$ , and wherein  $R^3$  is an acid amide or a free acid;

with the proviso that a maximum of one non-conservative amino acid exchange is effected per amino acid position in the sequence, wherein "non-conservative exchange" means an exchange of amino acids between the groups mentioned below:

Group I: Leu, Ile, Val, Met, His, Trp, Tyr, Phe,

Group II: Glu, Gln, Asp, Asn

Group III: Ser, Thr, Cys, Gly, Ala, Pro,

Group IV: Lys, Arg;

and

peptides selected from the group consisting of:

HWWRAESD-EARRSYNDPK-R<sup>2</sup>, HWYRATSDGEARRSYADPTSR<sup>2</sup>,

with the proviso that a maximum of two non-conservative amino acid exchanges are effected per amino acid position in the sequence, wherein "non-conservative exchange" means an exchange of amino acids between the groups mentioned below:

Group I: Leu, Ile, Val, Met, His, Trp, Tyr, Phe,

Group II: Glu, Gln, Asp, Asn,

Group III: Ser, Thr, Cys, Gly, Ala, Pro,

Group IV: Lys, Arg.

16. The peptides according to claim 13, characterized by being:

TGSFF SELWT SGKK-amide or free acid,

E YGSFF SELWT SGKK-amide or free acid,

T YGTLF SDFWL SGKK-amide or free acid,

His-Trp-Trp-Arg-Ala-Glu-Ser-Asp-Glu-Ala-Arg-Arg-Ser-Tyr-Asn-Asp-Pro-Lys-amide or free acid,

Ala-Arg-Arg-Cys-Tyr-Asn-Asp-Pro-Lys-amide or free acid,

D WGTLV SGFWE Y amide or free acid,

D WGTLF SDFWQ TGKK amide or free acid,

H WYRAT SDGEA RRSYA DPTSG KK-amide or free acid,

HWWRAESDEARRSYNDPKC-amide or free acid,

which may also be acetylated N-terminally.

- 17. The peptides according to claim 13, characterized by being bound by antibodies of patients suffering from dilatative cardiomyopathy.
- 18. The peptides according to claim 13, characterized in that said linker is selected from the group consisting of:
  - $\alpha$ -aminocarboxylic acids and their homo- and heterooligomers;
  - $\alpha,\omega$ -aminocarboxylic acids and their branched homo- or heterooligomers;
  - other amino acids and their linear and branched homo- or heterooligomers (peptides);
  - amino-oligoalkoxy-alkylamines;
  - maleinimidocarboxylic acid derivatives;
  - oligomers of alkylamines;
  - 4-alkylphenyl derivatives;
  - 4-oligoalkoxyphenyl or 4-oligoalkoxyphenoxy derivatives;
  - 4-oligoalkylmercaptophenyl or 4-oligoalkylmercaptophenoxy derivatives;

- 4-oligoalkylaminophenyl or 4-oligoalkylaminophenoxy derivatives;
- (oligoalkylbenzyl)phenyl or (4-oligoalkylbenzyl)phenoxy derivatives, and (4-oligoalkoxybenzyl)phenyl or (4-oligoalkoxybenzyl)phenoxy derivatives;
- trityl derivatives;
- benzyloxyaryl or benzyloxyalkyl derivatives;
- xanthene-3-yloxyalkyl derivatives;
- (4-alkylphenyl) or ω-(4-alkylphenoxy)alkanoic acid derivatives;
- oligoalkylphenoxyalkyl or oligoalkoxyphenoxyalkyl derivatives;
- carbamate derivatives;
- amines;
- trialkylsilyl or dialkylalkoxysilyl derivatives;
- alkyl or aryl derivatives;
- and combinations thereof.
- 19. The peptides according to claim 13, characterized by being bound to a solid phase.
- 20. The peptides according to claim 13, characterized by being bound to a solid phase through a spacer.
- 21. A medicament containing the peptides according to claim 13.
- 22. Use of the peptides according to claim 13 for the preparation of a medicament for treatment with diseases related to  $\beta_1$ -adrenergically active auto-antibodies, especially dilatative cardiomyopathy.
- 23. A method for treating diseases related to  $\beta_1$ -adrenergically active autoantibodies by removing the auto-antibodies by means of peptides according to claim 18 bound to a solid phase.
- 24. A device for chromatography containing peptides according to claim 18 bound to a solid phase.

#### **REMARKS**

The foregoing Preliminary Amendment is requested in order to delete the multiple dependent claims and avoid paying the multiple dependent claims fee.

Early action on the merits is respectfully requested.

Respectfully submitted,

JACOBSON HOLMAN PLLC

Βv

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Atty. Docket: P67710US0 Date: March 21, 2002

WEP:jrc

SMB

#### Peptides Against Auto-Antibodies Causing DCM

The present invention relates to peptides against auto-antibodies causing DCM, medicaments containing such peptides, the use of the peptides, methods for the treatment of diseases related to  $\beta_1$ -adrenergically active auto-antibodies, and a device for immunoadsorption containing the peptides bound to a solid phase.

The immune system is an essential component of all animal beings. In mammals, in particular, it serves for defense against microorganisms, tissue regeneration and destruction of tumor cells. In classical immunology, distinction is made between cellular and humoral immune defense. This means two distinguishable, but cooperating systems which ultimately represent the immune system.

A number of diseases exist which, due to their pathogenesis, are considered auto-immune diseases. In such diseases, the immune system of the afflicted subjects is directed against their own organs, tissues, cells or proteins and other molecules. The predominantly cell-mediated auto-immune diseases include multiple sclerosis and diabetes (type I).

A second group are the predominantly antibody-mediated auto-immune diseases. These include, for example, rheumatism, the less frequently occurring auto-immune diseases, such as myasthenia gravis or lupus erythematodes, and recently also dilatative cardiomyopathy (DCM).

The pathogenesis of most auto-immune diseases is unknown. There are various hypotheses and models of how to explain the genesis of auto-immune diseases. One explaining model is antigenic/molecular mimicry. In this model, it is

considered that microorganisms, e.g., viruses or parasites, provide themselves with particular molecules which are, for example, remarkably similar to or even in part identical with endogenous structures of the host and are therefore not recognized by the immune system of the host.

However, when they are recognized as foreign and antibodies are produced against them, then such antibodies will also recognize similar endogenous structures, which results in activation of the immune system and complement system. This induces pathological reactions in situ in the tissue, for example, chronic inflammations, or a pathological dysfunction of the cells occurs to which the auto-antibodies have bound.

Dilatative cardiomyopathy can be considered a prominent example thereof. In this auto-immune disease, the organisms erroneously forms antibodies which bind to defined regions of  $\beta_1$ -adrenergic receptor. These regions are on the first and second loops of a total of three extracellular loops of  $\beta_1$ -adrenergic receptor.

Such auto-antibodies which are capable to bind to these regions cause an increase of the pulsation rate in biological tests with rat cardiomyocytes in a cell culture (these cells have a nearly identical  $\beta_1$ -adrenergic receptor on their surface). This is referred to as a pharmaco-active effect of the auto-antibodies similar to that of adrenalin. The auto-antibodies directed against the epitopes on loops 1 and 2 of the  $\beta_1$ -adrenergic receptor are mainly observed in patients suffering from DCM. Occasionally, such auto-antibodies are also observed in patients having cardiac dysrhythmia and myocarditis.

Dilatative cardiomyopathy is an auto-immune disease which, when not treated, results in a severe deterioration of the cardiac output, i.e., reduction of the pumping output with simultaneous expansion of the myocardiac tissue by infiltrates, and then in heart transplantation or death.

However, if the antibodies are removed from the patient's blood by lavage of the blood, regeneration of the heart muscle and a dramatic improvement of the

myocardiac output, which almost reaches the values of healthy people, occur within one year.

In patients with DCM, an immunoglobulin fraction which contains the specific auto-antibodies binding to  $\beta_1$ -adrenergic receptor and thereby activate the cell can be isolated from the plasma. When peptides of  $\beta_1$ -adrenergic receptor which represent the binding site for the auto-antibodies are added to a cell culture of rat cardiomyocytes, the pathological effect of the immunoglobulin fraction can be neutralized.

If the same peptides which correspond to the native sequences are coupled to a solid phase, they are no longer capable of binding and eliminating the auto-antibodies described from a patient's blood plasma. This means that the peptides which correspond to the native sequence of  $\beta_1$ -adrenergic receptor and represent binding sites for the pathological auto-antibodies described cannot be used for immuno-adsorption.

It has been the object of the invention to provide peptides which recognize, bind and eliminate pathological auto-antibodies directed against functional epitopes in the blood or plasma of patients having a positive antibody state or DCM, wherein the peptides, in addition to the epitopes which respectively neutralize the antibody effect, simultaneously contain amino acid sequences which enable binding of the pathological antibodies.

Surprisingly, this object is achieved by peptides having the amino acid sequence

X01-X02-X03-G-X04-X05-X06-X07-X08-X09-W-X10-X11-X12

wherein

X01 = amino group, acetyl group, biotin group, fluorescent label, spacer, linker or deletion;

X02 = D,G,E,T,S or deletion;

X03 = W,Y,F,G,T;

X04 = T,S,A,G;

X12 = A, D, N, S;

X13 = D, N;

X14 = E, P; X15 = R, K, T; X16 = S, T, C, M or a deletion; X17 = amide, the free acid, GKK, SGKK or a spacer.

In particular, peptides according to the invention having the following amino acid sequence are employed:

X01-X02-X03-G-X04-X05-X06-X07-X08-X09-W-X10-X11-X12

wherein

X01 = amino group, acetyl group, biotin group, fluorescent label, spacer, linker or deletion;

X02 = D,E,T, or deletion;

X03 = W,Y,T;

X04 = T,S;

X05 = L,F;

X06 = V,F;

X07 = S;

X08 = G,D,E;

X09 = F,L;

X10 = E,Q,T,L;

X11 = Y,T,S;

X12 = amid, the free acid, GKK, or a spacer;

and peptides having the amino acid sequence

X01-H-W-X03-R-A-X05-S-D-X08-E-A-R-R-S-Y-X12-D-P-X15-X16-X17

wherein

X01 = amino group, amino acid, peptide, acetyl group, biotin group, fluorescent label, spacer, linker or deletion;

```
X03 = Y, W;
X05 = T, E;
X08 = G, \text{ or a deletion};
X12 = A, N;
X15 = K, T;
X16 = S, \text{ or a deletion};
X17 = \text{ amide, the free acid, GKK, SGKK or a spacer.}
```

It is particularly preferred to employ the following peptides which are selected from the group consisting of

```
-TGSFFSELWTSR<sup>2</sup>,
EYGSFFSELWTSR<sup>2</sup>,
TYGTLFSDFWLSR<sup>2</sup>,
DWGTLVSGFWEYR<sup>2</sup>,
DWGTLFSDFWQTR<sup>2</sup>,
```

wherein  $R^2$  is an acid amide, a free acid or  $GKKR^3$ , and wherein  $R^3$  is an acid amide or a free acid;

with the proviso that a maximum of one non-conservative amino acid exchange is effected per amino acid position in the sequence, wherein "non-conservative exchange" means an exchange of amino acids between the groups mentioned below:

Group I: Leu, Ile, Val, Met, His, Trp, Tyr, Phe,

Group II: Glu, Gln, Asp, Asn,

Group III: Ser, Thr, Cys, Gly, Ala, Pro,

Group IV: Lys, Arg;

and

peptides selected from the group consisting of:

HWWRAESD-EARRSYNDPK-R<sup>2</sup>, HWYRATSDGEARRSYADPTSR<sup>2</sup>,

with the proviso that a maximum of two non-conservative amino acid exchanges are effected per amino acid position in the sequence, wherein "non-conservative exchange" means an exchange of amino acids between the groups mentioned below:

Group I: Leu, Ile, Val, Met, His, Trp, Tyr, Phe,

Group II: Glu, Gln, Asp, Asn,

Group III: Ser, Thr, Cys, Gly, Ala, Pro,

Group IV: Lys, Arg.

The following peptides are particularly preferred:

TGSFF SELWT SGKK-amide or free acid, (SEQ. ID. NO. 1)

E YGSFF SELWT SGKK-amide or free acid, (SEQ. ID. NO. 2)

T YGTLF SDFWL SGKK-amide or free acid, (SEQ. ID. NO. 3)

His-Trp-Trp-Arg-Ala-Glu-Ser-Asp-Glu-Ala-Arg-Arg-Ser-Tyr-Asn-Asp-Pro-Lys-amide or free acid, (SEQ. ID. NO. 4)

Ala-Arg-Arg-Cys-Tyr-Asn-Asp-Pro-Lys-amide or free acid, (SEQ. ID. NO. 5)

D WGTLV SGFWE Y amide or free acid, (SEQ. ID. NO. 6)

D WGTLF SDFWQ TGKK amide or free acid, (SEQ. ID. NO. 7)

H WYRAT SDGEA RRSYA DPTSG KK-amide or free acid, (SEQ. ID. NO. 8)

HWWRAESDEARRSYNDPKC-amide or free acid, (SEQ. ID. NO. 9)

which may also be acetylated N-terminally.

The skilled person knows that amino acid positions in peptides or proteins can be exchanged in a conservative manner without affecting the function. In the

present case, a "conservative" exchange means an exchange within the groups set forth below:

Group I: Leu, Ile, Val, Met, His, Trp, Tyr, Phe,

Group II: Glu, Gln, Asp, Asn,

Group III: Ser, Thr, Cys, Gly, Ala, Pro,

Group IV: Lys, Arg.

The peptides according to the invention are bound, in particular, by antibodies of patients suffering from dilatative cardiomyopathy.

As a linker according to the invention, all structures may be used which are available for that purpose unless adversely affecting the binding behavior of the peptides towards the antibodies. Usually, a linker is a chemical compound which provides at least one linking site (functional group) on a polymeric matrix which is otherwise free of functions.

The linking site serves for coupling a ligand or a spacer and matches the chemical properties of the ligand or spacer. Such a link is stable or cleavable depending on the type of linker molecule.

A ligand is usually a compound having some special property. According to the invention, the ligand is preferably a peptide which is capable of specifically binding an auto-antibody which has an adrenergic activity and is directed against the  $\beta_1$ -adrenergic receptor of the heart muscle.

According to the invention, the following linkers are preferably employed:

 $\alpha$ -aminocarboxylic acids and their homo- and heterooligomers,  $\alpha$ , $\omega$ -aminocarboxylic acids and their branched homo- or heterooligomers, other amino acids and their linear and branched homo- or heterooligomers (peptides); amino-oligoalkoxy-alkylamines; maleinimidocarboxylic acid derivatives; oligomers of alkylamines; 4-alkylphenyl derivatives; 4-oligoalkoxyphenyl or 4-oligoalkoxy-

phenoxy derivatives; 4-oligoalkylmercaptophenyl or 4-oligoalkylmercaptophenoxy derivatives; 4-oligoalkylaminophenyl or 4-oligoalkylaminophenoxy derivatives; (oligoalkylbenzyl)phenyl or (4-oligoalkylbenzyl)phenoxy derivatives, and (4-oligoalkoxybenzyl)phenyl or (4-oligoalkoxybenzyl)phenoxy derivatives; trityl derivatives; benzyloxyaryl or benzyloxyalkyl derivatives; xanthene-3-yloxyalkyl derivatives; (4-alkylphenyl) or  $\omega$ -(4-alkylphenoxy)alkanoic acid derivatives; oligoalkylphenoxyalkyl or oligoalkoxyphenoxyalkyl derivatives; carbamate derivatives; amines; trialkylsilyl or dialkylalkoxysilyl derivatives; alkyl or aryl derivatives, and combinations thereof.

In particular, the peptides according to the invention are bound to a solid phase for use. Preferably, the binding of the peptides to the solid phase is effected through a spacer. As the spacer, there may be used virtually all chemical compounds or groups suitable for such a function unless adversely affecting the binding behavior to such an extent that binding of the antibody with the peptide is prevented or substantially impaired.

A spacer is usually a compound which is inserted between a ligand and a linker if necessary and serves for positioning the ligand at a distance and in a spatial position appropriate for the binding of the auto-antibody. Spacers are molecules having at least two chemically active groups (functional groups), of which one group binds to the linker molecule, and at least one second functional group mediates binding to a ligand. By selecting the spacer, an increase of flexibility and an improvement of accessibility as well as an oriented arrangement of the ligands and increase of ligand density on the surface can be achieved depending on requirements.

Spacers include, for example,  $\omega$ -aminocarboxylic acids and their homo- and heterooligomers,  $\alpha, \omega$ -aminocarboxylic acids and their branched homo- or heterooligomers, other aminocarboxylic acids and their linear and branched homo- or heterooligomers, maleinimidocarboxylic acid derivatives, hydroxycarboxylic acid derivatives, dicarboxylic acid derivatives, diamine derivatives dihydroxyalkyl derivatives, and hydroxyalkylamine derivatives. Preferably, mono- or dioligomers of  $\beta$ -alanine or  $\omega$ -aminohexanoic acid and branched mono- or dioligomers

of lysine or ornithine are used. The technology by means of which peptides can be anchored to solid phases is per se known to the skilled person.

In another embodiment of the invention, the peptides according to the invention are employed as medicaments.

In this concept, peptides are particularly altered (e.g. by cyclization) so that they cannot be destroyed by serum proteases and will bind antibodies in solutions. In this way, in-vivo neutralization of the antibodies can take place by intravenously administering the correspondingly processed peptides. The peptides are to be considered as medicaments herein. Their development is directly derived from the peptides binding the antibodies, which are fixed to a column matrix.

The amount of peptides to be administered depends on their molecular weight (i.e., their size) and on the concentration of auto-antibodies which can be reached in the blood stream and other compartments. According to what is known today, the amounts of auto-antibodies are within the µg and ng ranges. A quantity of between 1 and 5 µg of peptide should be sufficient for binding the antibodies present and pass them to elimination as an immune complex in accordance with the natural clearance mechanisms. Subsequently, the dosages can be lower since only the newly produced antibodies must be eliminated.

In principle, other dosage forms may be suitable in this case too. When the corresponding galenic methods are used, absorption in the intestine of peptides binding  $\beta_1$ -adrenergic antibodies should be achievable. In this case, the dosages would preferably be higher by a factor of about 10 to 20.

The peptides according to the invention can be employed for the preparation of a medicament for treatment of diseases related to  $\beta_1$ -adrenergically active autoantibodies, especially dilatative cardiomyopathy, blood hypertension diseases, and a form of cardiomyopathy induced by *Trypanosoma cruzi*. In addition to dilatative cardiomyopathy, a number of diseases exist which can be classified as auto-immune diseases and are subject to similar pathomechanisms. Also for preeclampsia and certain forms of malignant blood hypertension, auto-antibodies

have been described which contribute to the hyperactivation of cells by stimulating their angiotensin receptor or  $\alpha_1$  receptor, and participate in the genesis of defined clinical pictures. The use of peptides for eliminating these specific auto-antibodies or for in-vivo neutralization of the auto-antibodies can be regarded by analogy with dilatative cardiomyopathy.

According to the invention, a method is claimed for the treatment of diseases related to  $\beta_1$ -adrenergically active auto-antibodies by removing the auto-antibodies using peptides bound to a solid phase. Advantageously, this may be effected with a device for chromatography according to the invention containing the peptides according to the invention bound to a solid phase.

The peptides are fixed to a solid phase, e.g., sepharose, in a closed sterile container which usually has a volume of between 5 and 250 ml. In this sterile space, the patients' blood plasma, from which the cells have previously been removed by a medical engineering apparatus, flows over or through the adsorption matrix, i.e., the peptide-coated sepharose surface. This results in the binding of the pathological auto-antibodies to the peptides which simulate regions of the  $\beta_1$ -adrenergic receptor. If suitable adsorption matrices are employed, the previous separation of the cells can be dispensed with.

The remaining components of the blood plasma or blood including all necessary and useful immunoglobulins then leave the column and are recirculated into the patient's blood stream. This device for extracorporeal therapy is state of the art as far as the non-specific elimination of plasma proteins or immunoglobulin is concerned.

The device according to the invention uses the peptides derived from the  $\beta_1$ -adrenergic receptor for removing the small but pathologically relevant auto-antibody fraction from the plasma.

After a certain amount of blood plasma has flown through the device, the blood plasma stream can be switched over to a second column which is technically identical with the first, while the first column is being regenerated. That means,

the loaded antibodies are separated from the peptide and discarded by using different rinsing and elution solutions, preferably physiological saline with and without additional buffering, for example, by phosphate, glycine or citrate, and a pH range of from pH 2 to pH 7.5. Subsequently, the thus regenerated column is again available for the binding and elimination of the pathological auto-antibodies from the patient's blood plasma. This double column principle has proven useful and is always used when regeneration of the column is required.

A second variant of the use of peptides for the elimination of pathological autoantibodies includes the use of disposable columns. In these columns, the solidphase matrix contains an amount of peptide sufficient that major portions of the auto-antibodies can be removed from the plasma in several hours of treatment. The advantage resides in the fact that the time-consuming and tedious regeneration of the adsorption matrix can be dispensed with.

A third variant of the treatment of DCM patients by the elimination of pathological antibodies from the blood plasma includes the use of columns in which a previous separation of plasma and blood cells is not required due to the design of the columns.

The use of columns requires technical devices which ensure a blood and plasma input and flow adequate for the treatment by using different flexible tubes, pumps, monitor screens and other monitoring systems.

#### Example

#### 1. Peptides

The following two peptides were immobilized on cross-linked agarose beads Sepharose 4B as the solid phase.

Peptide 1: TGSFFCELWTSGKK

Peptide 2: HWWRAESDEARRSYNDPKC

Sepharose CL4B served as the filling matrix. From the two peptide matrices and the filling matrix, an affinity chromatographic column was prepared, and its function, i.e., the removal of DCM-related auto-antibodies from human plasma, was tested with human plasma as a sample.

#### 2. Immobilization of the peptides

For immobilization on a solid phase, the peptides were dissolved in coupling buffer (0.5 M NaCl, 0.1 M NaHCO3, pH 8.3) to a concentration of 2 mg/ml, and mixed with washed and CNBr-preactivated Sepharose 4B. After completion of the coupling reaction, the peptide matrices were washed with coupling buffer, and the excess CNBr groups were inactivated.

The peptide loading of the matrices was determined photometrically by calculating the difference between the mass of peptide employed before the coupling and the mass of non-immobilized peptide after the coupling.

The matrix loading with peptide 1 was 2.0 mg of peptide/ml of matrix. The matrix loading with peptide 2 was 1.8 mg of peptide/ml of matrix.

To prepare the affinity chromatographic column, peptide matrix 1 and peptide matrix 2 were mixed at a ratio of 1:1, and this mixture was mixed with filling matrix at a ratio of 1:5. The total volume of the matrix for the affinity chromatographic column was 100 ml.

#### 3. Removal of DCM-related auto-antibodies from human plasma

The function of the above described affinity chromatographic column was tested on two persons afflicted with DCM and exhibiting a positive test for DCM-related auto-antibodies.

Prior to the application, the persons must be treated with anti-coagulants, such as heparin, hirudin or citrate.

Thus, from the recommended concentration range of from 1500 to 3000 units, a bolus of heparin of 2000 units was chosen for intravenous administration. During the application, from the recommended concentration range of from 250 to 750 units of heparin per hour, a dosage of 500 units of heparin per hour was chosen for intravenous administration.

The blood was passed into a separator in which the separation of the cellular components of the blood from the blood plasma was effected. The plasma of the persons testing positive for DCM-related auto-antibodies was passed over the affinity chromatographic column. The volume ratio of the affinity matrix to the plasma was between 1:6 and 1:10 for the affinity chromatographic column runs performed. All in all, 20 such purification cycles were performed, wherein the treated plasma was reinfused to the persons after each cycle together with the cellular blood components. The total number of purification cycles of 20 is derived from 4 purification cycles per day of application for a total of 5 days of application.

#### 4. Quantitative assay of the DCM-related auto-antibodies

A quantitative assay of the DCM-related auto-antibodies was effected from plasma samples of the persons afflicted with DCM, obtained before and after the application on each day of application. The assay of the DCM-related auto-antibodies (antibodies against  $\beta_1$ -adrenergic receptor) was performed according to Wallukat, G., Wollenberger, A., Morwinski, R. and Pitschner, H.F. (1995); Anti- $\beta_1$ -adrenoceptor antibodies with chronotropic activity from the serum of patients with dilated cardiomyopathy: mapping of epitopes in the first and second extracellular loops; J. Mol. Cell. Cardiol. 27, 397-406.

# 5. Results of the removal of DCM-related Auto-Antibodies from Human Plasma

For person 1 having auto-antibodies which preferably bind to peptide 1, a reduction of auto-antibodies to 12% of the initial value was achieved after completion of the overall application.

For person 2 having auto-antibodies which preferably bind to peptide 2, a reduction of auto-antibodies to 5% of the initial value was achieved after completion of the overall application.

The results for the overall application are set forth in Table 1.

In both persons, the concentrations of other plasma parameters, such as total protein, albumin and IgG, were almost unaffected during the overall application.

Table 1: DCM-related auto-antibodies (antibodies against  $\beta_1$ -adrenergic receptor) in the human plasma of person 1 and person 2 before, during and after treatment with the affinity chromatographic column.

After elution from the affinity matrix, the auto-antibodies bound to the affinity chromatographic column were tested for preferential binding to peptide 1 or peptide 2 by means of peptide 1 and peptide 2 bound to a solid phase. Auto-antibody type 1 prefers the binding of peptide 1, and auto-antibody type 2 prefers the binding of peptide 2.

Person 1

Plasma sample	Auto-antibody 2	Auto-antibody 2
	[relative units]	[%]
before cycle 1-4	5.8	100
after cycle 1-4	2.4	41
before cycle 5-8	3.4	59
after cycle 5-8	1.6	28
before cycle 9-12	3.3	57
after cycle 9-12	2.0	34
before cycle 13-16	1.9	33
after cycle 13-16	0.8	14
before cycle 17-20	1.7	29
after cycle 17-20	0.7	12

Person 2

Plasma sample	Auto-antibody 2	Auto-antibody 2
	[relative units]	[%]
before cycle 1-4	6.1	100
after cycle 1-4	3.1	51
before cycle 5-8	3.8	62
after cycle 5-8	3.0	49
before cycle 9-12	3.5	57
after cycle 9-12	2.3	38
before cycle 13-16	2.8	46
after cycle 13-16	0.9	15
before cycle 17-20	1.7	28
after cycle 17-20	0.3	5

#### CLAIMS:

#### 1. A peptide having the amino acid sequence

X01-X02-X03-G-X04-X05-X06-X07-X08-X09-W-X10-X11-X12

#### wherein

X01 = amino group, acetyl group, biotin group, fluorescent label, spacer, linker or deletion;

X02 = D,G,E,T,S or deletion;

X03 = W,Y,F,G,T;

X04 = T,S,A,G;

X05 = L,F,Y,W;

X06 = V,I,W,F,Y;

X07 = S,A,C;

X08 = G,D,E,N,Q;

X09 = F,L,I,Y;

X10 = E,Q,T,S,L;

X11 = Y,F,T,S,W;

X12 = amide, the free acid, GKK, or a spacer;

and peptides having the amino acid sequence

X01-X02-W-X03-R-X04-X05-X06-X07-X08-E-A-R-X09-X10-X11-X12-X13-X14-X15-X16-X17

#### wherein

X01 = amino group, amino acid, peptide, acetyl group, biotin group, fluorescent label, spacer, linker or deletion;

X02 = H, E, Q;

X03 = H, F, Y, W; X04 = A, V; X05 = G, T, E, S, D, N; X06 = S, H, A; X07 = D, N, Q, E; X08 = G, A, or a deletion; X09 = D, N, R; X10 = S, T, C, M; X11 = H F, W, Y; X12 = A, D, N, S; X13 = D, N; X14 = E, P; X15 = R, K, T; X16 = S, T, C, M or a deletion; X17 = amide, the free acid, GKK, SGKK or a spacer.

2. The peptide according to claim 1 having the following amino acid sequence:

wherein

X01 = amino group, acetyl group, biotin group, fluorescent label, spacer, linker or deletion;

X02 = D,E,T, or deletion;

X03 = W,Y,T;

X04 = T,S;

X05 = L,F;

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X06 = V,F;
X07 = S;
X08 = G,D,E;
X09 = F_{,L};
X10 = E,Q,T,L;
X11 = Y,T,S;
X12 = amid, the free acid, GKK, or a spacer;
and peptides having the amino acid sequence
X01-H-W-X03-R-A-X05-S-D-X08-E-A-R-R-S-Y-X12-D-P-X15-X16-X17
wherein
X01 = amino group, amino acid, peptide, acetyl group, biotin group, fluo-
rescent label, spacer, linker or deletion;
X03 = Y, W;
X05 = T, E;
X08 = G, or a deletion;
X12 = A, N;
X15 = K, T;
X16 = S, or a deletion;
X17 = amide, the free acid, GKK, SGKK or a spacer.
The peptides according to any of claims 1 or 2 selected from the group
consisting of:
```

3.

-TGSFFSELWTSR<sup>2</sup>, EYGSFFSELWTSR<sup>2</sup>, TYGTLFSDFWLSR<sup>2</sup>, DWGTLVSGFWEYR<sup>2</sup>, DWGTLFSDFWQTR<sup>2</sup>,

wherein  $R^2$  is an acid amide, a free acid or  $GKKR^3$ , and wherein  $R^3$  is an acid amide or a free acid;

with the proviso that a maximum of one non-conservative amino acid exchange is effected per amino acid position in the sequence, wherein "nonconservative exchange" means an exchange of amino acids between the groups mentioned below:

Group I: Leu, Ile, Val, Met, His, Trp, Tyr, Phe,

Group II: Glu, Gln, Asp, Asn,

Group III: Ser, Thr, Cys, Gly, Ala, Pro,

Group IV: Lys, Arg;

and

peptides selected from the group consisting of:

HWWRAESD-EARRSYNDPK-R<sup>2</sup>, HWYRATSDGEARRSYADPTSR<sup>2</sup>,

with the proviso that a maximum of two non-conservative amino acid exchanges are effected per amino acid position in the sequence, wherein "non-conservative exchange" means an exchange of amino acids between the groups mentioned below:

Group I: Leu, Ile, Val, Met, His, Trp, Tyr, Phe,

Group II: Glu, Gln, Asp, Asn,

Group III: Ser, Thr, Cys, Gly, Ala, Pro,

Group IV: Lys, Arg.

4. The peptides according to claims 1 to 3, characterized by being:

TGSFF SELWT SGKK-amide or free acid,

E YGSFF SELWT SGKK-amide or free acid,

T YGTLF SDFWL SGKK-amide or free acid,

His-Trp-Trp-Arg-Ala-Glu-Ser-Asp-Glu-Ala-Arg-Arg-Ser-Tyr-Asn-Asp-Pro-Lys-amide or free acid,

Ala-Arg-Arg-Cys-Tyr-Asn-Asp-Pro-Lys-amide or free acid,

D WGTLV SGFWE Y amide or free acid,

D WGTLF SDFWQ TGKK amide or free acid,

H WYRAT SDGEA RRSYA DPTSG KK-amide or free acid,

HWWRAESDEARRSYNDPKC-amide or free acid,

which may also be acetylated N-terminally.

- 5. The peptides according to claims 1 to 4, characterized by being bound by antibodies of patients suffering from dilatative cardiomyopathy.
- 6. The peptides according to any of claims 1 to 5, characterized in that said linker is selected from the group consisting of:
  - α-aminocarboxylic acids and their homo- and heterooligomers;
  - $\alpha, \omega$ -aminocarboxylic acids and their branched homo- or heterooligomers;
  - other amino acids and their linear and branched homo- or heterooligomers (peptides);
  - amino-oligoalkoxy-alkylamines;
  - maleinimidocarboxylic acid derivatives;
  - oligomers of alkylamines;
  - 4-alkylphenyl derivatives;
  - 4-oligoalkoxyphenyl or 4-oligoalkoxyphenoxy derivatives;
  - 4-oligoalkylmercaptophenyl or 4-oligoalkylmercaptophenoxy derivatives;

- 4-oligoalkylaminophenyl or 4-oligoalkylaminophenoxy derivatives;
- (oligoalkylbenzyl)phenyl or (4-oligoalkylbenzyl)phenoxy derivatives, and (4-oligoalkoxybenzyl)phenyl or (4-oligoalkoxybenzyl)phenoxy derivatives;
- trityl derivatives;
- benzyloxyaryl or benzyloxyalkyl derivatives;
- xanthene-3-yloxyalkyl derivatives;
- (4-alkylphenyl) or ω-(4-alkylphenoxy)alkanoic acid derivatives;
- oligoalkylphenoxyalkyl or oligoalkoxyphenoxyalkyl derivatives;
- carbamate derivatives;
- amines;
- trialkylsilyl or dialkylalkoxysilyl derivatives;
- alkyl or aryl derivatives;
- and combinations thereof.
- 7. The peptides according to any of claims 1 to 6, characterized by being bound to a solid phase.
- 8. The peptides according to any of claims 1 to 7, characterized by being bound to a solid phase through a spacer.
- 9. A medicament containing the peptides according to any of claims 1 to 8.
- 10. Use of the peptides according to any of claims 1 to 8 for the preparation of a medicament for treatment with diseases related to  $\beta_1$ -adrenergically active auto-antibodies, especially dilatative cardiomyopathy.
- 11. A method for treating diseases related to  $\beta_1$ -adrenergically active autoantibodies by removing the auto-antibodies by means of peptides according to claim 6 or 7 bound to a solid phase.
- 12. A device for chromatography containing peptides according to claim 6 or 7 bound to a solid phase.

- 23 -

#### **Abstract**

Peptides which bind auto-antibodies being in a causally pathological relationship with dilatative cardiomyopathy are described. The peptides may be bound, for example, to a solid phase. Auto-antibodies can be removed by treating blood of patients suffering from DCM with the peptides according to the invention.

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